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 S  P  G  P  T  F  N  V  P  R  G  T  E  T  V  V  R  F  I  N  N  A  T  V  E  N  S  V  H  L    148

CACGGCTCCCCATCGCGTGCCCCTTTCGATGGTTGGGCTGAAGATGTGACCTTCCCTGGCGAGTACAAGGATTACTACTTTCCCAACTAC   540
 H  G  S  P  S  R  A  P  F  D  G  W  A  E  D  V  T  F  P  G  E  Y  K  D  Y  Y  F  P  N  Y    178

CAATCCGCCCGCCTTCTGTGGTACCATGACCACGCTTTCATGAAGACTGCTGAGAATGCCTACTTTGGTCAGGCTGGCGCCTACATTATC   630
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GATGGTACCCTGCGTTCGACCGAGGGTGAGGACCAGGACCTGTGGGGAGATGTCATCCATGTCAACGGACAGCCATGGCCTTTCCTTAAC   810
 D  G  T  L  R  S  T  E  G  E  D  Q  D  L  W  G  D  V  I  H  V  N  G  Q  P  W  P  F  L  N    268

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GCCCGAGTGCAGGAGCTGGCCGAGCAGGAGCCGTACAACCGCCTCGATGAGATCCTGGAGGATCTTGGAATCGAGGAGTAA          1791
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Figure 1

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Figure 2

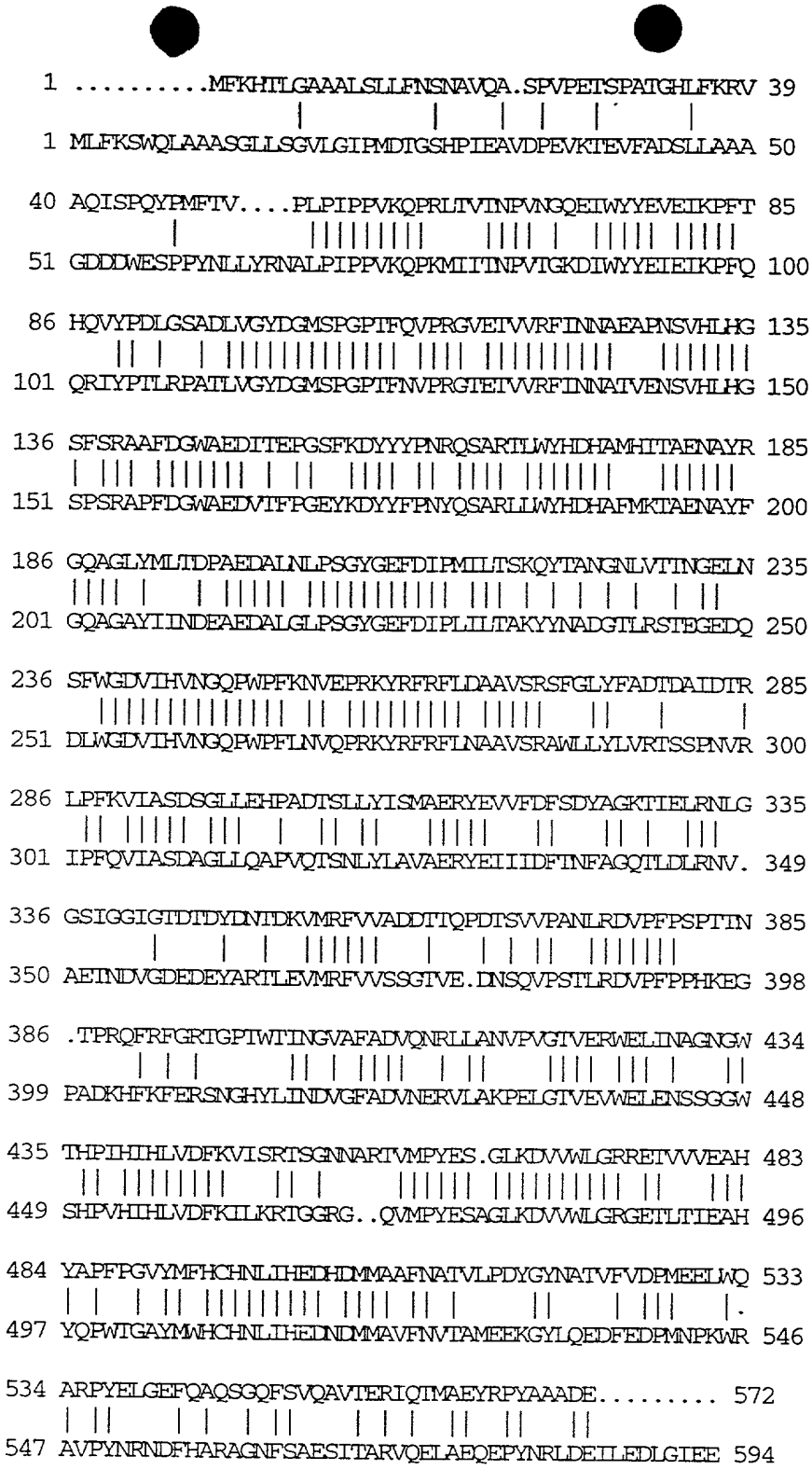
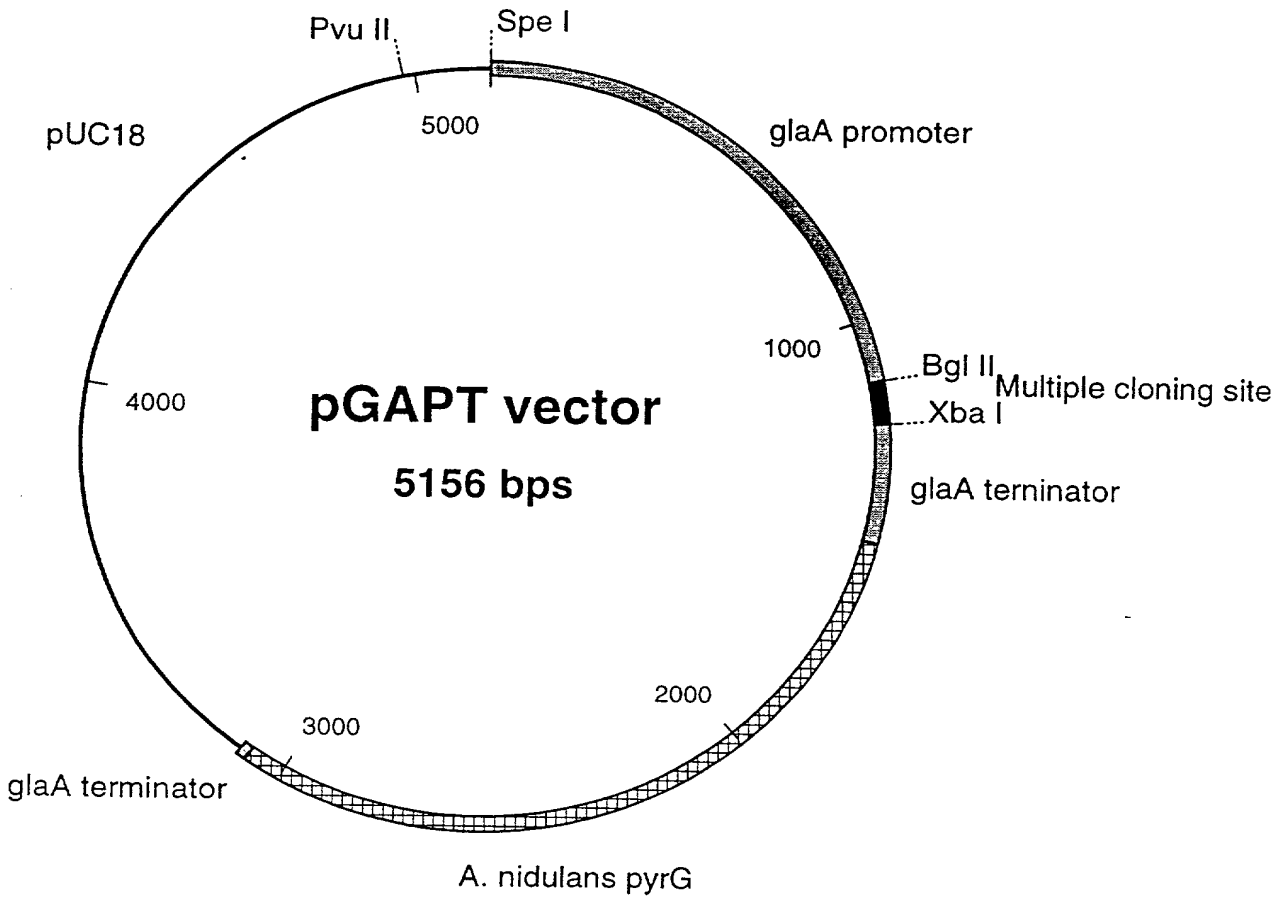


Figure 3: protein sequneces aligment of Bilirubin oxidase (top sequence) with Stachybotrys oxidase (bottom sequence).

Figure 3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2000
3000
glaA terminator
A. nidulans pyrG

Figure 4

AGATCTAATA TGCTGTTCAA GTCATGGCAA CTGGCAGCAG CCTCCGGGCT CCTGTCTGGA 60
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CCTCCATACA ACTTGCTTTA CAGGTGAGAC ACCTGTCCCA CCTGTTTTCC CTCGATAACT 240
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Figure 5

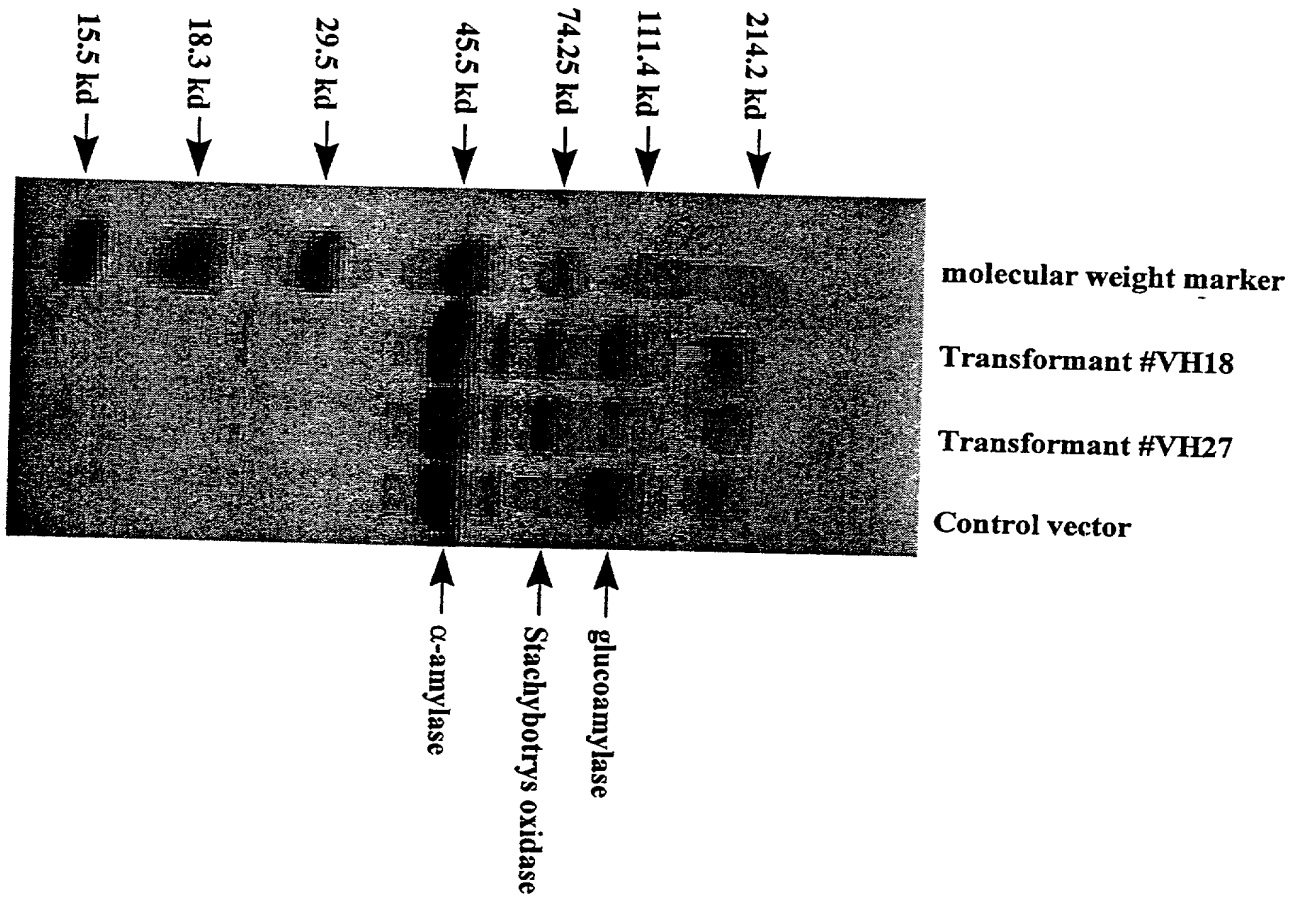
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Figure 6